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                        GenCore version 5.1.3
              Copyright (c) 1993 - 2003 Compugen Ltd.

OM protein - nucleic search, using frame_plus_p2n model

Run on:          January 16, 2003, 17:00:17 ; Search time 50.5143 Seconds
                                              (without alignments)
                                              114.746 Million cell updates/sec

Title:           US-09-856-070-19
Perfect score:   65
Sequence:        1 KEELMERLQDYEE 13

Scoring table:   BLOSUM62
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                 Ygapop 10.0 , Ygapext  0.5
                 Fgapop  6.0 , Fgapext  7.0
                 Delop   6.0 , Delext   7.0

Searched:        393868 seqs, 222934149 residues

Total number of hits satisfying chosen parameters:      787736

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Command line parameters:
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-DB=Published_Applications_NA -QFMT=fastap -SUFFIX=rnpb -MINMATCH=0.1
-LOOPCL=0 -LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62
-TRANS=human40.cdi -LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100
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-MAXLEN=2000000000 -USER=US09856070_@CGN_1_1_87_@runat_14012003_155835_1681
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-DEV_TIMEOUT=120 -WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6
-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7

Database :       Published_Applications_NA:*
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                 4:  /cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*
                 5:  /cgn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*
                 6:  /cgn2_6/ptodata/2/pubpna/PCTUS_PUBCOMB.seq:*
                 7:  /cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*
                 8:  /cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*
                 9:  /cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*
                 10:  /cgn2_6/ptodata/2/pubpna/US09_PUBCOMB.seq:*
                 11:  /cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*
                 12:  /cgn2_6/ptodata/2/pubpna/US10_PUBCOMB.seq:*
                 13:  /cgn2_6/ptodata/2/pubpna/US60_NEW_PUB.seq:*
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     Pred. No. is the number of results predicted by chance to have a
     score greater than or equal to the score of the result being printed,
     and is derived by analysis of the total score distribution.

                              SUMMARIES
                    %
Result              Query
  No.     Score     Match Length DB  ID                          Description
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    1        65     100.0   2930  10  US-09-960-253-156          Sequence 156, App
    2        65     100.0   3044  10  US-09-880-107-3718         Sequence 3718, Ap
    3        65     100.0   3047  10  US-09-864-864-329          Sequence 329, App
    4        65     100.0   3115  10  US-09-925-299-123          Sequence 123, App
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c   5     40    61.5     224  10  US-09-864-761-29393         Sequence 29393, A
c   6     40    61.5     520  10  US-09-864-761-12828         Sequence 12828, A
    7     40    61.5    2166  12  US-10-025-187-1             Sequence 1, Appli
    8     40    61.5    2170   9  US-09-854-133-728          Sequence 728, App
    9     40    61.5   45839  12  US-10-025-187-3             Sequence 3, Appli
c  10     39    60.0     205  10  US-09-864-761-27935         Sequence 27935, A
c  11     39    60.0     452  10  US-09-864-761-11355         Sequence 11355, A
   12     39    60.0     550   9  US-10-046-935-2209         Sequence 2209, Ap
   13     39    60.0     550   9  US-09-878-178-2209         Sequence 2209, Ap
   14     39    60.0    1488  10  US-09-863-475A-9            Sequence 9, Appli
   15     39    60.0    3647  10  US-09-863-475A-7            Sequence 7, Appli
   16     39    60.0    4242  12  US-10-044-090-190           Sequence 190, App
   17     38    58.5    1134  10  US-09-873-676-29            Sequence 29, Appl
c  18     38    58.5    2124   9  US-09-938-842A-854         Sequence 854, App
   19     38    58.5    2302   9  US-09-764-868-445          Sequence 445, App
   20     38    58.5    2313   9  US-09-854-133-157          Sequence 157, App
   21     38    58.5    2313  10  US-09-738-973-157           Sequence 157, App
   22     38    58.5    2314   9  US-09-764-868-12           Sequence 12, Appl
   23     38    58.5    2497  10  US-09-946-893-1             Sequence 1, Appli
   24     38    58.5   16086  10  US-09-764-877-2385          Sequence 2385, Ap
   25     37    56.9     210   9  US-09-796-692-7066         Sequence 7066, Ap
c  26     37    56.9     591  10  US-09-864-761-9272          Sequence 9272, Ap
   27     37    56.9     629  10  US-09-864-761-31941         Sequence 31941, A
   28     37    56.9    1780  10  US-09-917-800A-1521         Sequence 1521, Ap
   29     37    56.9    1960  10  US-09-746-801A-12           Sequence 12, Appl
   30     37    56.9    3617  10  US-09-746-801A-14           Sequence 14, Appl
   31     37    56.9    7420  10  US-09-917-800A-502          Sequence 502, App
   32     37    56.9    7787  10  US-09-954-456-2006          Sequence 2006, Ap
   33   36.5    56.2    1653  10  US-09-965-703-66            Sequence 66, Appl
   34   36.5    56.2    1705  10  US-09-965-697-13            Sequence 13, Appl
   35   36.5    56.2    1798  10  US-09-956-998A-16           Sequence 16, Appl
   36   36.5    56.2    5620   9  US-10-125-751-21           Sequence 21, Appl
   37   36.5    56.2    5620   9  US-09-912-552-21           Sequence 21, Appl
   38   36.5    56.2    5620  10  US-09-918-029-21            Sequence 21, Appl
   39   36.5    56.2    5620  12  US-10-038-271-21            Sequence 21, Appl
c  40   36.5    56.2    7312  10  US-09-946-334-12            Sequence 12, Appl
   41   36.5    56.2    7460   9  US-10-038-001-7            Sequence 7, Appli
c  42   36.5    56.2    7788  10  US-09-946-334-11            Sequence 11, Appl
   43   36.5    56.2    8937  10  US-09-872-733-8             Sequence 8, Appli
   44   36.5    56.2    8937  10  US-09-872-733-9             Sequence 9, Appli
c  45   36.5    56.2   14683   9  US-10-109-853-1            Sequence 1, Appli


                              ALIGNMENTS

RESULT 1
US-09-960-253-156
; Sequence 156, Application US/09960253
; Patent No. US20020123619A1
; GENERAL INFORMATION:
;  APPLICANT: Benson, Darin R.
;  APPLICANT:  Mohamath, Raodoh
;  APPLICANT:  Lodes, Michael J.
;  TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
;  TITLE OF INVENTION:  AND DIAGNOSIS OF LUNG CANCER
;  FILE REFERENCE: 210121.556
;  CURRENT APPLICATION NUMBER: US/09/960,253
;  CURRENT FILING DATE:  2001-09-20
;  NUMBER OF SEQ ID NOS: 187
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 156
;   LENGTH: 2930
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-960-253-156

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Percent Similarity:       100.00%        Conservative:    0
Best Local Similarity:    100.00%        Mismatches:      0
Query Match:              100.00%        Indels:          0
DB:                       10             Gaps:            0
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US-09-856-070-19 (1-13) x US-09-960-253-156 (1-2930)

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Db     1106 AAGGAGGAGTTGATGCTGCGGCTGCAGGACTATGAGGAG 1144

RESULT 2
US-09-880-107-3718
; Sequence 3718, Application US/09880107
; Patent No. US20020142981A1
; GENERAL INFORMATION:
;  APPLICANT: Horne, Darci T.
;  APPLICANT:  Vockley, Joseph G.
;  APPLICANT:  Scherf, Uwe
;  APPLICANT:  Gene Logic, Inc.
;  TITLE OF INVENTION: Gene Expression Profiles in Liver Cancer
;  FILE REFERENCE: 44921-5028-WO
;  CURRENT APPLICATION NUMBER: US/09/880,107
;  CURRENT FILING DATE:  2001-06-14
;  PRIOR APPLICATION NUMBER: US 60/211,379
;  PRIOR FILING DATE: 2000-06-14
;  PRIOR APPLICATION NUMBER: US 60/237,054
;  PRIOR FILING DATE: 2000-10-02
;  NUMBER OF SEQ ID NOS: 3950
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 3718
;    LENGTH: 3044
;    TYPE: DNA
;    ORGANISM: Homo sapiens
;    FEATURE:
;    OTHER INFORMATION: Genbank Accession No. US20020142981A1 X51521
;    NAME/KEY: unsure
;    LOCATION: (1)..(3044)
;    OTHER INFORMATION: n - a or c or g or t
US-09-880-107-3718

Alignment Scores:
Pred. No.:                 0.000429       Length:         3044
Score:                     65.00          Matches:        13
Percent Similarity:        100.00%        Conservative:   0
Best Local Similarity:     100.00%        Mismatches:     0
Query Match:               100.00%        Indels:         0
DB:                        10             Gaps:           0

US-09-856-070-19 (1-13) x US-09-880-107-3718 (1-3044)

Qy        1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
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Db     1147 AAGGAGGAGTTGATGCTGCGGCTGCAGGACTATGAGGAG 1185

RESULT 3
US-09-864-864-329
; Sequence 329, Application US/09864864
; Patent No. US20020102679A1
; GENERAL INFORMATION:
;  APPLICANT: Xu, Jiangchun
;  APPLICANT:  Mitcham, Jennifer L.
;  APPLICANT:  Harlocker, Susan L.
;  APPLICANT:  Dillon, Davin C.
;  APPLICANT:  Secrist, Heather
;  APPLICANT:  Lodes, Michael J.
;  APPLICANT:  Algate, Paul A.
;  APPLICANT:  Fling, Steve P.
;  APPLICANT:  Mannion, Jane
;  APPLICANT:  Benson, Darin R.
;  APPLICANT:  Carter, Darrick
;  TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
;  TITLE OF INVENTION:  AND DIAGNOSIS OF OVARIAN CANCER
;  FILE REFERENCE: 210121.523
;  CURRENT APPLICATION NUMBER: US/09/864,864
;  CURRENT FILING DATE:  2001-05-23
;  NUMBER OF SEQ ID NOS: 341
;  SOFTWARE: Corixa Invention Disclosure Database
; SEQ ID NO 329
;    LENGTH: 3047
;    TYPE: DNA
;    ORGANISM: Homo sapiens
;    FEATURE:
;    NAME/KEY: misc_feature
;    LOCATION: (1)...(3047)
;    OTHER INFORMATION: n - A,T,C or G
US-09-864-864-329

Alignment Scores:
Pred. No.:                 0.000429       Length:         3047
Score:                     65.00          Matches:        13
Percent Similarity:        100.00%        Conservative:   0
Best Local Similarity:     100.00%        Mismatches:     0
Query Match:               100.00%        Indels:         0
DB:                        10             Gaps:           0

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Qy        1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
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Db     1147 AAGGAGGAGTTGATGCTGCGGCTGCAGGACTATGAGGAG 1185

RESULT 4
US-09-925-299-123
; Sequence 123, Application US/09925299
; Patent No. US20020055627A1
; GENERAL INFORMATION:
;  APPLICANT: Rosen et al.
;  TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
;  FILE REFERENCE: PA102
;  CURRENT APPLICATION NUMBER: US/09/925,299
;  CURRENT FILING DATE:  2001-08-10
;  PRIOR APPLICATION NUMBER: PCT/US00/05883
;  PRIOR FILING DATE: 2000-03-08
;  PRIOR APPLICATION NUMBER: 60/124,270
;  PRIOR FILING DATE: 1999-03-12
;  NUMBER OF SEQ ID NOS: 1556
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 123
;    LENGTH: 3115
;    TYPE: DNA
;    ORGANISM: Homo sapiens
US-09-925-299-123

Alignment Scores:
Pred. No.:                 0.00044        Length:         3115
Score:                     65.00          Matches:        13
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Best Local Similarity:     100.00%        Mismatches:     0
Query Match:               100.00%        Indels:         0
DB:                        10             Gaps:           0

US-09-856-070-19 (1-13) x US-09-925-299-123 (1-3115)

Qy        1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
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Db     1179 AAGGAGGAGTTGATGCTGCGGCTGCAGGACTATGAGGAG 1217

RESULT 5
US-09-864-761-29393/c
; Sequence 29393, Application US/09864761
; Patent No. US20020048763A1
; GENERAL INFORMATION:
;  APPLICANT: Penn, Sharron G.
;  APPLICANT:  Rank, David R.
;  APPLICANT:  Hanzel, David K.
;  APPLICANT:  Chen, Wensheng
;  TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
;  TITLE OF INVENTION:  GENE EXPRESSION ANALYSIS BY MICROARRAY
;  FILE REFERENCE: Aeomica-X-1
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;  CURRENT APPLICATION NUMBER: US/09/864,761
;  CURRENT FILING DATE:  2001-05-23
;  PRIOR APPLICATION NUMBER: US 60/180,312
;  PRIOR FILING DATE: 2000-02-04
;  PRIOR APPLICATION NUMBER: US 60/207,456
;  PRIOR FILING DATE: 2000-05-26
;  PRIOR APPLICATION NUMBER: US 09/632,366
;  PRIOR FILING DATE: 2000-08-03
;  PRIOR APPLICATION NUMBER: GB 24263.6
;  PRIOR FILING DATE: 2000-10-04
;  PRIOR APPLICATION NUMBER: US 60/236,359
;  PRIOR FILING DATE: 2000-09-27
;  PRIOR APPLICATION NUMBER: PCT/US01/00666
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00667
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00664
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00669
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00665
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00668
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00663
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00662
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00661
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00670
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: US 60/234,687
;  PRIOR FILING DATE: 2000-09-21
;  PRIOR APPLICATION NUMBER: US 09/608,408
;  PRIOR FILING DATE: 2000-06-30
;  PRIOR APPLICATION NUMBER: US 09/774,203
;  PRIOR FILING DATE: 2001-01-29
;  NUMBER OF SEQ ID NOS: 49117
;  SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
; SEQ ID NO 29393
;   LENGTH: 224
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: MAP TO AC009155.3
;   OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL - 0.65
;   OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL - 0.62
;   OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL - 0.98
;   OTHER INFORMATION: EST_HUMAN HIT: AL138321.1, EVALUE 1.00e-118
;   OTHER INFORMATION: SWISSPROT HIT: P28684, EVALUE 8.70e+00
;   OTHER INFORMATION: NT HIT: AF079406.1, EVALUE 1.30e-01
US-09-864-761-29393

Alignment Scores:
Pred. No.:                 2.98            Length:         224
Score:                     40.00           Matches:        7
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Best Local Similarity:     53.85%          Mismatches:     2
Query Match:               61.54%          Indels:         0
DB:                        10              Gaps:           0

US-09-856-070-19 (1-13) x US-09-864-761-29393 (1-224)

Qy        1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
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Db       57 AAGCAGAATCTGTTGCTGGAACTCCGTAACTATGAGGAA 19

RESULT 6
US-09-864-761-12828/c
; Sequence 12828, Application US/09864761
; Patent No. US20020048763A1
; GENERAL INFORMATION:
;  APPLICANT: Penn, Sharron G.
;  APPLICANT: Rank, David R.
;  APPLICANT: Hanzel, David K.
;  APPLICANT: Chen, Wensheng
;  TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
;  TITLE OF INVENTION:  GENE EXPRESSION ANALYSIS BY MICROARRAY
;  FILE REFERENCE: Aeomica-X-1
;  CURRENT APPLICATION NUMBER: US/09/864,761
;  CURRENT FILING DATE:  2001-05-23
;  PRIOR APPLICATION NUMBER: US 60/180,312
;  PRIOR FILING DATE: 2000-02-04
;  PRIOR APPLICATION NUMBER: US 60/207,456
;  PRIOR FILING DATE: 2000-05-26
;  PRIOR APPLICATION NUMBER: US 09/632,366
;  PRIOR FILING DATE: 2000-08-03
;  PRIOR APPLICATION NUMBER: GB 24263.6
;  PRIOR FILING DATE: 2000-10-04
;  PRIOR APPLICATION NUMBER: US 60/236,359
;  PRIOR FILING DATE: 2000-09-27
;  PRIOR APPLICATION NUMBER: PCT/US01/00666
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00667
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00664
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00669
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00665
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00668
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00663
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00662
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00661
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00670
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: US 60/234,687
;  PRIOR FILING DATE: 2000-09-21
;  PRIOR APPLICATION NUMBER: US 09/608,408
;  PRIOR FILING DATE: 2000-06-30
;  PRIOR APPLICATION NUMBER: US 09/774,203
;  PRIOR FILING DATE: 2001-01-29
;  NUMBER OF SEQ ID NOS: 49117
;  SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
; SEQ ID NO 12828
;   LENGTH: 520
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: MAP TO AC009155.3
;   OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL - 0.65
;   OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL - 0.62
;   OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL - 0.98
US-09-864-761-12828

Alignment Scores:
Pred. No.:                 7.94            Length:         520
Score:                     40.00           Matches:        7
Percent Similarity:        84.62%          Conservative:   4
Best Local Similarity:     53.85%          Mismatches:     2
Query Match:               61.54%          Indels:         0
DB:                        10              Gaps:           0

US-09-856-070-19 (1-13) x US-09-864-761-12828 (1-520)

Qy        1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
              |||:::   |||:::|||   |||::::::|||||||||
Db      150 AAGCAGAATCTGTTGCTGGAACTCCGTAACTATGAGGAA 112

RESULT 7
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US-10-025-187-1
; Sequence 1, Application US/10025187
; Patent No. US20020150931A1
; GENERAL INFORMATION:
;  APPLICANT: SHEFFIELD, VAL
;  APPLICANT:  NISHIMURA, DARRYL
;  APPLICANT:  STONE, EDWARD
;  TITLE OF INVENTION: A BARDET-BIEDL SUSCEPTIBILITY GENE AND USES THEREOF
;  FILE REFERENCE: IOWA:034US
;  CURRENT APPLICATION NUMBER: US/10/025,187
;  CURRENT FILING DATE:  2001-12-18
;  PRIOR APPLICATION NUMBER: 60/256,900
;  PRIOR FILING DATE: 2000-12-19
;  NUMBER OF SEQ ID NOS: 3
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
;   LENGTH: 2166
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-025-187-1

Alignment Scores:
Pred. No.:                41.9          Length:         2166
Score:                    40.00         Matches:        7
Percent Similarity:       84.62%        Conservative:   4
Best Local Similarity:    53.85%        Mismatches:     2
Query Match:              61.54%        Indels:         0
DB:                       12            Gaps:           0

US-09-856-070-19 (1-13) x US-10-025-187-1 (1-2166)

Qy          1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
                |||:::   |||:::|||   |||::::::|||||||||
Db       1033 AAGCAGAATCTGTTGCTGGAACTCCGTAACTATGAGGAA 1071

RESULT 8
US-09-854-133-728
; Sequence 728, Application US/09854133
; Publication No. US20020183499A1
; GENERAL INFORMATION:
;  APPLICANT: Lodes, Michael J.
;  APPLICANT:  Mohamath, Raodoh
;  APPLICANT:  Henderson, Robert A.
;  APPLICANT:  Benson, Darin R.
;  APPLICANT:  Secrist, Heather
;  TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
;  TITLE OF INVENTION:  THE THERAPY AND DIAGNOSIS OF LUNG CANCER
;  FILE REFERENCE: 210121.475C10
;  CURRENT APPLICATION NUMBER: US/09/854,133
;  CURRENT FILING DATE:  2001-05-11
;  NUMBER OF SEQ ID NOS: 735
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 728
;   LENGTH: 2170
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(2170)
;   OTHER INFORMATION: n = A,T,C or G
US-09-854-133-728

Alignment Scores:
Pred. No.:                42            Length:         2170
Score:                    40.00         Matches:        9
Percent Similarity:       76.92%        Conservative:   1
Best Local Similarity:    69.23%        Mismatches:     3
Query Match:              61.54%        Indels:         0
DB:                       9             Gaps:           0

US-09-856-070-19 (1-13) x US-09-854-133-728 (1-2170)

Qy          1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
              |||||||||||||||   ||||||:::      ||||||
Db       1208 AAGGAAGAGCTAATGGAACGTCTAAAACAAATTGAAGAG 1246

RESULT 9
US-10-025-187-3
; Sequence 3, Application US/10025187
; Patent No. US20020150931A1
; GENERAL INFORMATION:
;  APPLICANT: SHEFFIELD, VAL
;  APPLICANT:  NISHIMURA, DARRYL
;  APPLICANT:  STONE, EDWARD
;  TITLE OF INVENTION: A BARDET-BIEDL SUSCEPTIBILITY GENE AND USES THEREOF
;  FILE REFERENCE: IOWA:034US
;  CURRENT APPLICATION NUMBER: US/10/025,187
;  CURRENT FILING DATE:  2001-12-18
;  PRIOR APPLICATION NUMBER: 60/256,900
;  PRIOR FILING DATE: 2000-12-19
;  NUMBER OF SEQ ID NOS: 3
;  SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 3
;   LENGTH: 45839
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-025-187-3

Alignment Scores:
Pred. No.:                1.47e+03      Length:         45839
Score:                    40.00         Matches:        7
Percent Similarity:       84.62%        Conservative:   4
Best Local Similarity:    53.85%        Mismatches:     2
Query Match:              61.54%        Indels:         0
DB:                       12            Gaps:           0

US-09-856-070-19 (1-13) x US-10-025-187-3 (1-45839)

Qy          1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
              |||:::   |||:::|||   |||::::::|||||||||
Db      24930 AAGCAGAATCTGTTGCTGGAACTCCGTAACTATGAGGAA 24968

RESULT 10
US-09-864-761-27935/c
; Sequence 27935, Application US/09864761
; Patent No. US20020048763A1
; GENERAL INFORMATION:
;  APPLICANT: Penn, Sharron G.
;  APPLICANT:  Rank, David R.
;  APPLICANT:  Hanzel, David K.
;  APPLICANT.  Chen, Wensheng
;  TITLE OF INVENTION: HUMAN GENOME DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
;  TITLE OF INVENTION.  GENE EXPRESSION ANALYSIS BY MICROARRAY
;  FILE REFERENCE: Aeomica-X-1
;  CURRENT APPLICATION NUMBER: US/09/864,761
;  CURRENT FILING DATE:  2001-05-23
;  PRIOR APPLICATION NUMBER: US 60/180,312
;  PRIOR FILING DATE: 2000-02-04
;  PRIOR APPLICATION NUMBER: US 60/207,456
;  PRIOR FILING DATE: 2000-05-26
;  PRIOR APPLICATION NUMBER: US 09/632,366
;  PRIOR FILING DATE: 2000-08-03
;  PRIOR APPLICATION NUMBER: GB 24263.6
;  PRIOR FILING DATE: 2000-10-04
;  PRIOR APPLICATION NUMBER: US 60/236,359
;  PRIOR FILING DATE: 2000-09-27
;  PRIOR APPLICATION NUMBER: PCT/US01/00666
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00667
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00664
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00669
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00665
;  PRIOR FILING DATE: 2001-01-30
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;  PRIOR APPLICATION NUMBER: PCT/US01/00668
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00663
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00662
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00661
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00670
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: US 60/234,687
;  PRIOR FILING DATE: 2000-09-21
;  PRIOR APPLICATION NUMBER: US 09/608,408
;  PRIOR FILING DATE: 2000-06-30
;  PRIOR APPLICATION NUMBER: US 09/774,203
;  PRIOR FILING DATE: 2001-01-29
;  NUMBER OF SEQ ID NOS: 49117
;  SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
; SEQ ID NO 27935
;   LENGTH: 205
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: MAP TO AC006195.1
;   OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL - 1.2
;   OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL - 1.2
;   OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL - 7.6
;   OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL - 1.4
;   OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL - 1.1
;   OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL - 1.1
;   OTHER INFORMATION: SWISSPROT HIT: P38110, EVALUE 1.70e+00
;   OTHER INFORMATION: NT HIT: AF095771.1, EVALUE 6.00e-93
;   OTHER INFORMATION: EST_HUMAN HIT: AA453960.1, EVALUE 5.00e-88
US-09-864-761-27935

Alignment Scores:
Pred. No.:                4.32          Length:        205
Score:                    39.00         Matches:       8
Percent Similarity:       90.91%        Conservative:  2
Best Local Similarity:    72.73%        Mismatches:    1
Query Match:              60.00%        Indels:        0
DB:                       10            Gaps:          0

US-09-856-070-19 (1-13) x US-09-864-761-27935 (1-205)

Qy        3 GluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
            ||||||:::|||||||||||||:::|||   |||
Db      151 GAGCTTATTCTTCGCCTTCAAGAATATTTTGAA 119

RESULT 11
US-09-864-761-11355/c
; Sequence 11355, Application US/09864761
; Patent No. US20020048763A1
; GENERAL INFORMATION:
;  APPLICANT: Penn, Sharron G.
;  APPLICANT:  Rank, David R.
;  APPLICANT:  Hanzel, David K.
;  APPLICANT:  Chen, Wensheng
;  TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
;  TITLE OF INVENTION:  GENE EXPRESSION ANALYSIS BY MICROARRAY
;  FILE REFERENCE: Aeomica-X-1
;  CURRENT APPLICATION NUMBER: US/09/864,761
;  CURRENT FILING DATE:  2001-05-23
;  PRIOR APPLICATION NUMBER: US 60/180,312
;  PRIOR FILING DATE: 2000-02-04
;  PRIOR APPLICATION NUMBER: US 60/207,456
;  PRIOR FILING DATE: 2000-05-26
;  PRIOR APPLICATION NUMBER: US 09/632,366
;  PRIOR FILING DATE: 2000-08-03
;  PRIOR APPLICATION NUMBER: GB 24263.6
;  PRIOR FILING DATE: 2000-10-04
;  PRIOR APPLICATION NUMBER: US 60/236,359
;  PRIOR FILING DATE: 2000-09-27
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;  PRIOR APPLICATION NUMBER: PCT/US01/00666
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00667
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00664
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00669
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00665
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00668
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00663
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00662
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00661
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: PCT/US01/00670
;  PRIOR FILING DATE: 2001-01-30
;  PRIOR APPLICATION NUMBER: US 60/234,687
;  PRIOR FILING DATE: 2000-09-21
;  PRIOR APPLICATION NUMBER: US 09/608,408
;  PRIOR FILING DATE: 2000-06-30
;  PRIOR APPLICATION NUMBER: US 09/774,203
;  PRIOR FILING DATE: 2001-01-29
;  NUMBER OF SEQ ID NOS: 49117
;  SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
; SEQ ID NO 11355
;   LENGTH: 452
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   OTHER INFORMATION: MAP TO AC006195.1
;   OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL - 1.2
;   OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1.2
;   OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 7.6
;   OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 1.4
;   OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL - 1.1
;   OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 1.1
US-09-864-761-11355

Alignment Scores:
Pred. No.:                10.9          Length:        452
Score:                    39.00         Matches:       8
Percent Similarity:       90.91%        Conservative:  2
Best Local Similarity:    72.73%        Mismatches:    1
Query Match:              60.00%        Indels:        0
DB:                       10            Gaps:          0

US-09-856-070-19 (1-13) x US-09-864-761-11355 (1-452)

Qy        3 GluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
            ||||||:::|||||||||||||:::|||   |||
Db      396 GAGCTTATTCTTCGCCTTCAAGAATATTTTGAA 364

RESULT 12
US-10-046-935-2209
; Sequence 2209, Application US/10046935
; Patent No. US20020156011A1
; GENERAL INFORMATION:
;  APPLICANT: Jiang, Yuqiu
;  APPLICANT:  Harlocker, Susan L.
;  APPLICANT:  Secrist, Heather
;  APPLICANT:  Wang, Aijun
;  APPLICANT:  Stolk, John A.
;  TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
;  TITLE OF INVENTION:  AND DIAGNOSIS OF COLON CANCER
;  FILE REFERENCE: 210121.527C1
;  CURRENT APPLICATION NUMBER: US/10/046,935
;  CURRENT FILING DATE:  2002-01-15
;  NUMBER OF SEQ ID NOS: 2239
;  SOFTWARE: FastSEQ for Windows Version 4.0
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; SEQ ID NO 2209
;   LENGTH: 550
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: 12, 472, 488, 495
;   OTHER INFORMATION: n - A,T,C or G
US-10-046-935-2209

Alignment Scores:
Pred. No.:               13.6          Length:        550
Score:                   39.00         Matches:       7
Percent Similarity:      84.62%        Conservative:  4
Best Local Similarity:   53.85%        Mismatches:    2
Query Match:             60.00%        Indels:        0
DB:                      9             Gaps:          0

US-09-856-070-19 (1-13) x US 10 046 935 2209 (1 550)

Qy        1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
              |||   |||::::::|||:::||||||:::|||   |||
Db      286 AAATCTCACATACTTTTAAAATTACAACAATATTTACAA 324

RESULT 13
US 09-878-178-2209
; Sequence 2209, Application US/09878178
; Patent No. US20020177552A1
; GENERAL INFORMATION:
;  APPLICANT: Jiang, Yuqiu
;  APPLICANT:  Harlocker, Susan L.
;  APPLICANT:  Secrist, Heather
;  TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
;  TITLE OF INVENTION:  AND DIAGNOSIS OF COLON CANCER
;  FILE REFERENCE: 210121.527
;  CURRENT APPLICATION NUMBER: US/09/878,178
;  CURRENT FILING DATE:  2001-06-08
;  NUMBER OF SEQ ID NOS: 2237
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 2209
;   LENGTH: 550
;   TYPE: DNA
;   ORGANISM: Homo sapien
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(550)
;   OTHER INFORMATION: n - A,T,C or G
US-09-878-178-2209

Alignment Scores:
Pred. No.:               13.6          Length:        550
Score:                   39.00         Matches:       7
Percent Similarity:      84.62%        Conservative:  4
Best Local Similarity:   53.85%        Mismatches:    2
Query Match:             60.00%        Indels:        0
DB:                      9             Gaps:          0

US-09-856-070-19 (1-13) x US-09-878-178-2209 (1-550)

Qy        1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
              |||   |||......|||...||||||:..|||   |||
Db      286 AAATCTGAGATAGTTTTAAAATTACAAGAATATTTAGAA 324

RESULT 14
US-09-863-475A-9
; Sequence 9, Application US/09863475A
; Patent No. US20020102688A1
;   GENERAL INFORMATION:
;         APPLICANT: LOWE, JOHN B.
;         TITLE OF INVENTION: METHODS AND PRODUCTS FOR THE SYNTHESIS
;                             OF OLIGOSACCHARIDE STRUCTURES ON GLYCOPROTEINS,
;                             GLYCOLIPIDS, OR AS FREE MOLECULES, AND FOR THE ISOLATION
;                             OF CLONED GENETIC SEQUENCES THAT DETERMINE THESE STRUCTURES
;         NUMBER OF SEQUENCES: 14
;         CORRESPONDENCE ADDRESS:
;              ADDRESSEE: OBLON, SPIVAK, McCLELLAND, MAIER & NEUSTADT,
;                         P.C.
;              STREET: 1755 Jefferson Davis Highway, Fourth Floor
;              CITY: Arlington
;              STATE: Virginia
;              COUNTRY: U.S.A.
;              ZIP: 22202
;         COMPUTER READABLE FORM:
;              MEDIUM TYPE: Floppy disk
;              COMPUTER: IBM PC compatible
;              OPERATING SYSTEM: PC-DOS/MS-DOS
;              SOFTWARE: PatentIn Release #1.0, Version #1.25
;         CURRENT APPLICATION DATA:
;              APPLICATION NUMBER: US/09/863,475A
;              FILING DATE: 24-May-2001
;              CLASSIFICATION: <Unknown>
;         PRIOR APPLICATION DATA:
;              APPLICATION NUMBER: 07/914,281
;              FILING DATE: 20-JUL-1992
;         ATTORNEY/AGENT INFORMATION:
;              NAME: Lavalleye, Jean-Paul M. P.
;              REGISTRATION NUMBER: 31,451
;              REFERENCE/DOCKET NUMBER: 2363-060-55
;         TELECOMMUNICATION INFORMATION:
;              TELEPHONE: (703)521-4500
;              TELEFAX: (703)486-2347
;              TELEX: 248855 OPAT UR
;    INFORMATION FOR SEQ ID NO: 9:
;         SEQUENCE CHARACTERISTICS:
;              LENGTH: 1488 base pairs
;              TYPE: nucleic acid
;              STRANDEDNESS: unknown
;              TOPOLOGY: unknown
;         MOLECULE TYPE: DNA (genomic)
;         SEQUENCE DESCRIPTION: SEQ ID NO: 9:
US-09-863-475A-9

Alignment Scores:
Pred. No.:               43.5          Length:        1488
Score:                   39.00         Matches:       8
Percent Similarity:      83.33%        Conservative:  2
Best Local Similarity:   66.67%        Mismatches:    2
Query Match:             60.00%        Indels:        0
DB:                      10            Gaps:          0

US-09-856-070-19 (1-13) x US-09-863-475A-9 (1-1488)

Qy        2 GluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
              ||||||:::   ||||||:::   ||||||||||||
Db      679 GAGGAGGTGGATCTGCGCGTGTTGGACTACGAGGAG 714

RESULT 15
US-09-863-475A-7
; Sequence 7, Application US/09863475A
; Patent No. US20020102688A1
;   GENERAL INFORMATION:
;         APPLICANT: LOWE, JOHN B.
;         TITLE OF INVENTION: METHODS AND PRODUCTS FOR THE SYNTHESIS
;                             OF OLIGOSACCHARIDE STRUCTURES ON GLYCOPROTEINS,
;                             GLYCOLIPIDS, OR AS FREE MOLECULES, AND FOR THE ISOLATION
;                             OF CLONED GENETIC SEQUENCES THAT DETERMINE THESE STRUCTU
;         NUMBER OF SEQUENCES: 14
;         CORRESPONDENCE ADDRESS:
;              ADDRESSEE: OBLON, SPIVAK, McCLELLAND, MAIER & NEUSTADT,
;                         P.C.
;              STREET: 1755 Jefferson Davis Highway, Fourth Floor
;              CITY: Arlington
;              STATE: Virginia
;              COUNTRY: U.S.A.
;              ZIP: 22202
;         COMPUTER READABLE FORM:
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;             MEDIUM TYPE: Floppy disk
;             COMPUTER: IBM PC compatible
;             OPERATING SYSTEM: PC-DOS/MS-DOS
;             SOFTWARE: PatentIn Release #1.0, Version #1.25
;        CURRENT APPLICATION DATA:
;             APPLICATION NUMBER: US/09/863,475A
;             FILING DATE: 24-May-2001
;             CLASSIFICATION: <Unknown>
;        PRIOR APPLICATION DATA:
;             APPLICATION NUMBER: 07/914,281
;             FILING DATE: 20-JUL-1992
;        ATTORNEY/AGENT INFORMATION:
;             NAME: Lavalleye, Jean-Paul M. P.
;             REGISTRATION NUMBER: 31,451
;             REFERENCE/DOCKET NUMBER: 2363-060-55
;        TELECOMMUNICATION INFORMATION:
;             TELEPHONE: (703)521-4500
;             TELEFAX: (703)486-2347
;             TELEX: 248855 OPAT UR
;   INFORMATION FOR SEQ ID NO: 7:
;        SEQUENCE CHARACTERISTICS:
;             LENGTH: 3647 base pairs
;             TYPE: nucleic acid
;             STRANDEDNESS: unknown
;             TOPOLOGY: unknown
;        MOLECULE TYPE: DNA (genomic)
;        ANTI-SENSE: NO
;        SEQUENCE DESCRIPTION: SEQ ID NO: 7:
US-09-863-475A-7

Alignment Scores:
Pred. No.:                 124           Length:        3647
Score:                     39.00         Matches:       8
Percent Similarity:        83.33%        Conservative:  2
Best Local Similarity:     66.67%        Mismatches:    2
Query Match:               60.00%        Indels:        0
DB:                        10            Gaps:          0

US-09-856-070-19 (1-13) x US-09-863-475A-7 (1-3647)

Qy        2 GluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
            ||||||:::   ||||||:::   ||||||||||||
Db     2347 GAGGAGGTGGATCTGCGCGTGTTGGACTACGAGGAG 2382


Search completed: January 16, 2003, 21.46:08
Job time : 55.5143 secs
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